



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/669,162A
Source: IFWO
Date Processed by STIC: 5/7/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/669,162A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 ☐ Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 ☐ Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 ☐ Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 ☐ Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 ☒ Variable Length Sequence(s) 6, 7, 8, 9, maybe more contain h's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 ☐ PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 ☐ Skipped Sequences
 (OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 ☐ Skipped Sequences
 (NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 ☐ Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 ☐ Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 ☐ Use of <220> Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 ☐ PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 ☐ Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



IFWU

RAW SEQUENCE LISTING

DATE: 05/07/2004

PATENT APPLICATION: US/10/669,162A

TIME: 14:10:39

Input Set : A:\250060016U2.txt

Output Set: N:\CRF4\05072004\J669162A.raw

3 <110> APPLICANT: Breaker, Ronald R.
 4 Nahvi, Ali
 5 Sudarsan, Narasimhan
 6 Ebert, Margaret S.
 7 Winkler, Wade
 8 Barrick, Jeffrey E.
 9 Wickiser, John K.
 11 <120> TITLE OF INVENTION: RIBOSWITCHES, METHODS FOR THEIR USE, AND
 12 COMPOSITIONS FOR USE WITH RIBOSWITCHES
 14 <130> FILE REFERENCE: 25006.0016U2
 16 <140> CURRENT APPLICATION NUMBER: 10/669,162A
 17 <141> CURRENT FILING DATE: 2003-09-22
 19 <150> PRIOR APPLICATION NUMBER: 60/412,468
 20 <151> PRIOR FILING DATE: 2002-09-20
 22 <160> NUMBER OF SEQ ID NOS: 377
 24 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 202
 28 <212> TYPE: RNA
 29 <213> ORGANISM: Escherichia coli
 31 <400> SEQUENCE: 1
 32 gccgguccug ugaguuuaa gggaauccag ugcgaaucug gagcugacgc gcagcgguaa 60
 33 ggaaaggugc gaugauugcg uuaugcggac acugccauuc gguggggaagu caucaucucu 120
 34 uaguaucuua gauacccuc caagcccgaa gaccugccgg ccaacgucgc aucugguucu 180
 35 caucaucgcg uaauauugau ga 202
 37 <210> SEQ ID NO: 2
 38 <211> LENGTH: 165
 39 <212> TYPE: RNA
 40 <213> ORGANISM: Escherichia coli
 42 <220> FEATURE:
 43 <221> NAME/KEY: misc_feature
 44 <222> LOCATION: 155
 45 <223> OTHER INFORMATION: r = a or g
 47 <220> FEATURE:
 48 <221> NAME/KEY: misc_feature
 49 <222> LOCATION: 157
 50 <223> OTHER INFORMATION: y = c or (t/u)
 52 <400> SEQUENCE: 2
 53 ggaaccaaac gacucggggu gcccucugc gugaaggcug agaaauaccc guaucaccug 60
 54 aucuggauaa ugccagcgua gggagucac ggaccaccag gucauugcuu cuucacguua 120
 55 uggcaggagc aaacuaugca agucgaccug cuggruycag cgcaa 165
 57 <210> SEQ ID NO: 3
 58 <211> LENGTH: 240

pp 1-7

u only. '+' cannot appear in an RNA sequence.

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TIME: 14:10:39

Input Set : A:\250060016U2.txt

Output Set: N:\CRF4\05072004\J669162A.raw

```

59 <212> TYPE: RNA
60 <213> ORGANISM: Escherichia coli
62 <220> FEATURE:
63 <221> NAME/KEY: misc_feature
64 <222> LOCATION: (155)...(240)
65 <223> OTHER INFORMATION: n = g, a, c, or t/u
67 <400> SEQUENCE: 3
68 ggaaugcccc auuugcgggg cuauuuucuu gucggagugc cuuaacuggc ugagaccguu 60
69 uauucgggau ccgcggaacc ugaucaggcu aaauaccugcg aagggaacaa gaguuaaucu 120
-> 70 gcuaucgcau cgccccugcg gcgaucgucu cuugnnnnnn nnnnnnnnnn nnnnnnnnnn 180
71 nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 240
73 <210> SEQ ID NO: 4
74 <211> LENGTH: 165
75 <212> TYPE: RNA
76 <213> ORGANISM: Escherichia coli
78 <220> FEATURE:
79 <221> NAME/KEY: misc_feature
80 <222> LOCATION: 65, 74, 107, 130
81 <223> OTHER INFORMATION: s = g or c
83 <220> FEATURE:
84 <221> NAME/KEY: misc_feature
85 <222> LOCATION: 25, 26, 34, 35, 64, 75, 106, 131
86 <223> OTHER INFORMATION: w = a or t/u
88 <400> SEQUENCE: 4
89 ggaaccaaac gacucggggu gcccwucugc gugwwggcug agaaauaccc guaucaccug 60
90 aucwsgauaa ugcswgcgua gggaagucac ggaccaccag gucauwscuu cuucacguua 120
91 uggcaggags waacuaugca agucgaccug cuggauccag cgcaa 165
93 <210> SEQ ID NO: 5
94 <211> LENGTH: 176
95 <212> TYPE: RNA
96 <213> ORGANISM: Artificial Sequence
98 <220> FEATURE:
99 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
100     synthetic construct
102 <220> FEATURE:
103 <221> NAME/KEY: misc_feature
104 <222> LOCATION: (39)...(156)
105 <223> OTHER INFORMATION: n = g, a, c or t/u
107 <400> SEQUENCE: 5
-> 108 ggauaauagc cguagguugc gaaagcgacc cugaguagnn nnnnncaaga gaagcagagg 60
109 gacuggcccg acgaagcuuc agcaaccggu guaauggcga ucagccauga ccaaggugcu 120
110 aaauccagca agcucgaaca gcuuggaagn nnnnnncgaa acgguagcga gaggcuc 176
113 <210> SEQ ID NO: 6
114 <211> LENGTH: 97
115 <212> TYPE: RNA
116 <213> ORGANISM: Artificial Sequence
118 <220> FEATURE:
119 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
120     synthetic construct

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TIME: 14:10:39

Input Set : A:\250060016U2.txt

Output Set: N:\CRF4\05072004\J669162A.raw

122 <220> FEATURE:
 123 <221> NAME/KEY: misc_feature
 124 <222> LOCATION: 1, 6, 26, 58, 66, 76, 97
 125 <223> OTHER INFORMATION: n = a variable number of any nucleotide
 127 <220> FEATURE:
 128 <221> NAME/KEY: misc_feature
 129 <222> LOCATION: 5, 7, 8, 11, 12, 18-20, 24, 25, 29, 30, 33-35, 38, 40, 41,
 130 47, 50, 54-56, 59, 60, 75, 77-79, 85, 89, 93
 131 <223> OTHER INFORMATION: n = g, a, c or t/u u
 133 <220> FEATURE:
 134 <221> NAME/KEY: misc_feature
 135 <222> LOCATION: 27, 36, 48, 53, 57, 80, 87
 136 <223> OTHER INFORMATION: r = a or g
 138 <220> FEATURE:
 139 <221> NAME/KEY: misc_feature
 140 <222> LOCATION: 67, 83
 141 <223> OTHER INFORMATION: y = c or t/u u
 143 <400> SEQUENCE: 6
 -> 144 nggunnnnaa nngggaannn ggunnnrann cennnrngn ncccgcnrcn guinnnninn 60
 145 cacugnyggg aaggnnnnr agynggrana ccngcn 97
 147 <210> SEQ ID NO: 7
 148 <211> LENGTH: 56
 149 <212> TYPE: RNA
 150 <213> ORGANISM: Artificial Sequence
 152 <220> FEATURE:
 153 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
 154 synthetic construct
 156 <220> FEATURE:
 157 <221> NAME/KEY: misc_feature u
 158 <222> LOCATION: 7, 50
 159 <223> OTHER INFORMATION: d = g, a or t/u (u)
 161 <220> FEATURE:
 162 <221> NAME/KEY: misc_feature
 163 <222> LOCATION: 1, 8, 13, 36, 56
 164 <223> OTHER INFORMATION: n = a variable number of any nucleotide
 166 <220> FEATURE:
 167 <221> NAME/KEY: misc_feature
 168 <222> LOCATION: 2-5, 17-20, 21-24, 30-34, 38-40, 41-43, 45-47
 169 <223> OTHER INFORMATION: n = g, a, c or t/u u
 171 <220> FEATURE:
 172 <221> NAME/KEY: misc_feature
 173 <222> LOCATION: 54
 174 <223> OTHER INFORMATION: r = a or g
 177 <400> SEQUENCE: 7
 -> 178 nnnngdncu gaganannnn nnnnaccugn nnnncunnnn nnnngnnncgd aggran 56
 180 <210> SEQ ID NO: 8
 181 <211> LENGTH: 97
 182 <212> TYPE: RNA
 183 <213> ORGANISM: Artificial Sequence

variable length not permitted
 (see item 5 on
 Error
 Summary
 sheet)

see item 5 on Error Summary
 sheet

RAW SEQUENCE LISTING

DATE: 05/07/2004

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TIME: 14:10:39

Input Set : A:\250060016U2.txt

Output Set: N:\CRF4\05072004\J669162A.raw

185 <220> FEATURE:

186 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =

187 synthetic construct

189 <220> FEATURE:

190 <221> NAME/KEY: misc_feature

191 <222> LOCATION: 57, 62

192 <223> OTHER INFORMATION: k = g or (t/u) u

194 <220> FEATURE:

195 <221> NAME/KEY: misc_feature

196 <222> LOCATION: 37, 47

197 <223> OTHER INFORMATION: n = a variable number of any nucleotide

199 <220> FEATURE:

200 <221> NAME/KEY: misc_feature

201 <222> LOCATION: 11, 17, 20, 28, 36, 46, 48, 58, 61, 77-79

202 <223> OTHER INFORMATION: n = g, a, c or (t/u) u

204 <220> FEATURE:

205 <221> NAME/KEY: misc_feature

206 <222> LOCATION: 6, 35, 43, 54, 59, 65-68, 74, 90, 91, 95-97

207 <223> OTHER INFORMATION: r = a or g

209 <220> FEATURE:

210 <221> NAME/KEY: misc_feature

211 <222> LOCATION: 1-3, 15, 31, 40, 44, 51-53, 64, 84

212 <223> OTHER INFORMATION: y = c or (t/u) u

214 <400> SEQUENCE: 8

-> 215 **yyyucrgggc nggggynaau ucccnaccgg yggurnnag yccrygnnga yyyrguknra** 60

216 **nkcyrrrrcc gacrgunnna gucyggaugr ragarr** 97

218 <210> SEQ ID NO: 9

219 <211> LENGTH: 86

220 <212> TYPE: (RNA)

221 <213> ORGANISM: Artificial Sequence

223 <220> FEATURE:

224 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =

225 synthetic construct

227 <220> FEATURE:

228 <221> NAME/KEY: misc_feature

229 <222> LOCATION: 52, 72

230 <223> OTHER INFORMATION: n = a variable number of any nucleotide

232 <220> FEATURE:

233 <221> NAME/KEY: misc_feature

234 <222> LOCATION: 1, 7-9, 13, 14, 16, 18, 25, 26, 32, 33, 37, 39, 42, 43, 50,

235 51, 53-55, 62, 63, 66-69, 71, 73, 75, 76, 78, 79, 86

236 <223> OTHER INFORMATION: n = g, a, c or (t/u) u

238 <220> FEATURE:

239 <221> NAME/KEY: misc_feature

240 <222> LOCATION: 38, 44, 70, 77, 83

241 <223> OTHER INFORMATION: r = a or g

243 <220> FEATURE:

244 <221> NAME/KEY: misc_feature

245 <222> LOCATION: 17, 34, 60, 74

RAW SEQUENCE LISTING

DATE: 05/07/2004

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TIME: 14:10:39

Input Set : A:\250060016U2.txt

Output Set: N:\CRF4\05072004\J669162A.raw

246 <223> OTHER INFORMATION: y = c or (t/u) M

248 <400> SEQUENCE: 9

-> 249 ncuuaunnnng agnngnynga gggannggcc cnnyganinc cnnergcaacn nnnnngugcy 60
250 annccnnnnr nnnynnrnng auragn 86

252 <210> SEQ ID NO: 10

253 <211> LENGTH: 69

254 <212> TYPE: (RNA)

255 <213> ORGANISM: Artificial Sequence

257 <220> FEATURE:

258 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
259 synthetic construct

261 <220> FEATURE:

262 <221> NAME/KEY: misc_feature

263 <222> LOCATION: 1, 2, 10-17, 22, 25-31, 34, 40-46, 54-60, 68, 69

264 <223> OTHER INFORMATION: n = g, a, c or (t/u) M

266 <220> FEATURE:

267 <221> NAME/KEY: misc_feature

268 <222> LOCATION: 5, 18, 67

269 <223> OTHER INFORMATION: r = a or g

271 <220> FEATURE:

272 <221> NAME/KEY: misc_feature

273 <222> LOCATION: 65

274 <223> OTHER INFORMATION: y = c or (t/u) M

276 <400> SEQUENCE: 10

-> 277 nnucruauan nnnnnnnrau anggnnnnnn ngunucuacn nnnnnnccgu aaannnnnnn 60
278 acuaygrnn 69

280 <210> SEQ ID NO: 11

281 <211> LENGTH: 69

282 <212> TYPE: (RNA)

283 <213> ORGANISM: Artificial Sequence

285 <220> FEATURE:

286 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
287 synthetic construct

289 <220> FEATURE:

290 <221> NAME/KEY: misc_feature

291 <222> LOCATION: 1, 2, 10-17, 22, 25-31, 34, 40-46, 54-60, 68, 69

292 <223> OTHER INFORMATION: n = g, a, c or (t/u) M

294 <220> FEATURE:

295 <221> NAME/KEY: misc_feature

296 <222> LOCATION: 5, 18, 67

297 <223> OTHER INFORMATION: r = a or g

299 <220> FEATURE:

300 <221> NAME/KEY: misc_feature

301 <222> LOCATION: 65

302 <223> OTHER INFORMATION: y = c or (t/u) M

304 <400> SEQUENCE: 11

-> 305 nnucruauan nnnnnnnrau anggnnnnnn ngunucuacn nnnnnnccgu aaannnnnnn 60
306 auuaygrnn 69

308 <210> SEQ ID NO: 12

FYI

The types of errors shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.

RAW SEQUENCE LISTING ERROR SUMMARY
 PATENT APPLICATION: US/10/669,162A

DATE: 05/07/2004
 TIME: 14:10:40

Input Set : A:\250060016U2.txt
 Output Set: N:\CRF4\05072004\J669162A.raw

Base Note:

g of n and/or Xaa have been detected in the Sequence Listing. Please review the
 yuence Listing to ensure that a corresponding explanation is presented in the <220>
 <223> fields of each sequence which presents at least one n or Xaa.

I#:3; N Pos. 155,156,157,158,159,160,161,162,163,164,165,166,167,168,169
 I#:3; N Pos. 170,171,172,173,174,175,176,177,178,179,180,181,182,183,184
 I#:3; N Pos. 185,186,187,188,189,190,191,192,193,194,195,196,197,198,199
 I#:3; N Pos. 200,201,202,203,204,205,206,207,208,209,210,211,212,213,214
 I#:3; N Pos. 215,216,217,218,219,220,221,222,223,224,225,226,227,228,229
 I#:3; N Pos. 230,231,232,233,234,235,236,237,238,239,240
 I#:5; N Pos. ~~39~~,40,41,42,43,44,45,150,151,152,153,154,155,156
 I#:6; N Pos. ~~1,5,6,7,8,11,12,18,19,20,24,25,26,29,30,33,34,35,38,40,41,47~~
 I#:6; N Pos. 50,54,55,56,58,59,60,66,75,76,77,78,79,85,89,93,97
 I#:7; N Pos. ~~1,2,3,4,5,8,15,17,18,19,20,21,22,23,24,30,31,32,33,34,36,38~~
 I#:7; N Pos. 39,40,41,42,43,45,46,47,56
 I#:8; N Pos. ~~11,17,20,25,36,37,46,47,48,58,61,77,78,79~~
 I#:9; N Pos. ~~1,7,8,9,18,1A,16,18,25,26,32,33,37,39,42,48,50,51,52,53,54~~
 I#:9; N Pos. 55,62,63,66,67,68,69,71,72,73,75,76,78,79,86
 I#:10; N Pos. ~~1,2,10,11,12,13,14,15,16,17,22,25,26,27,28,29,30,31,34,40~~
 I#:10; N Pos. 41,42,43,44,45,46,54,55,56,57,58,59,60,68,68
 I#:11; N Pos. ~~1,2,10,11,12,13,14,15,16,17,22,25,26,27,28,29,30,31,34,40~~
 I#:11; N Pos. 41,42,43,44,45,46,54,55,56,57,58,59,60,68,69
 I#:12; N Pos. 1,13,14,15,16,17,18,20,21,26,27,28,29,30,31,32,33,34,35,39
 I#:12; N Pos. 40,41,42,43,44,45,46,47,48,49,50,51,52,53,59,60,61,62,63,64
 I#:12; N Pos. 65,66,67,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93
 I#:12; N Pos. 94,95,96,97,98,99,100,101,107,108,109,110,111,112,113,114
 I#:12; N Pos. 115,116,117,118,121,122,123,124,125,126,127,128,129,130,131
 I#:12; N Pos. 132,139,145,151
 I#:15; N Pos. 33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52
 I#:15; N Pos. 53,54,55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72
 I#:15; N Pos. 73,74,75,76,77,78,79,80,81,82,83
 I#:16; N Pos. 33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52
 I#:16; N Pos. 53,54,55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72
 I#:16; N Pos. 73,74,75,76,77,78,79,80,81,82,83
 I#:17; N Pos. 15,16,17,18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34
 I#:17; N Pos. 35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54
 I#:17; N Pos. 55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74
 I#:17; N Pos. 75,76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93,94
 I#:17; N Pos. 95,96,97,98,99,100,101,102,103,104,105,106,107,108,109,110
 I#:17; N Pos. 111,112,113,114,115,116,117,118,119,120,121,122,123
 I#:18; N Pos. 1,2,3,4,5,6,7,10,11,12,13,14,15,16,23,25,26,27,28,29,30,31
 I#:18; N Pos. 32,40,41,42,44,46,47,48,49,50,51,53,54,55,64,65,66
 I#:19; N Pos. 12,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,42,43,44
 I#:19; N Pos. 45,46,47,48,49,50,51
 I#:20; N Pos. 12,20,21,22,23,24,25,26,27,28,29
 I#:21; N Pos. 12,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38
 I#:21; N Pos. 39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54,55,56,57,58
 I#:21; N Pos. 59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74,75,76,77,78

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/669,162A

DATE: 05/07/2004
TIME: 14:10:40

Input Set : A:\250060016U2.txt
Output Set: N:\CRF4\05072004\J669162A.raw

I#:21; N Pos. 79,89,94
I#:22; N Pos. 19,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38
I#:22; N Pos. 39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54,55,56,57,58
I#:22; N Pos. 59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74,75,76,77,78
I#:22; N Pos. 79,80,81,82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98
I#:22; N Pos. 99,100,101,102,103,104,105,106,107,108,109,110,111,112,113
I#:22; N Pos. 114,115,116,117,118,119,120,121,122,123,124,125,126,127,128

VERIFICATION SUMMARY

DATE: 05/07/2004

PATENT APPLICATION: US/10/669,162A

TIME: 14:10:40

Input Set : A:\250060016U2.txt

Output Set: N:\CRF4\05072004\J669162A.raw

70 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:120
341 Repeated in SeqNo=3
108 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
341 Repeated in SeqNo=5
144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
341 Repeated in SeqNo=6
178 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
215 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
341 Repeated in SeqNo=8
249 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
341 Repeated in SeqNo=9
277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
341 Repeated in SeqNo=10
305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
341 Repeated in SeqNo=11
349 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
341 Repeated in SeqNo=12
384 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
341 Repeated in SeqNo=15
399 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
341 Repeated in SeqNo=16
114 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
341 Repeated in SeqNo=17
144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
341 Repeated in SeqNo=18
158 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
186 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
341 Repeated in SeqNo=21
501 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
341 Repeated in SeqNo=22
517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
341 Repeated in SeqNo=23
533 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
341 Repeated in SeqNo=24
548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
341 Repeated in SeqNo=25
563 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
341 Repeated in SeqNo=26
578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
341 Repeated in SeqNo=27
594 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
341 Repeated in SeqNo=28
509 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
341 Repeated in SeqNo=29
525 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
341 Repeated in SeqNo=30
541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0

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341 Repeated in SeqNo=31

557 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0

1713 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:98